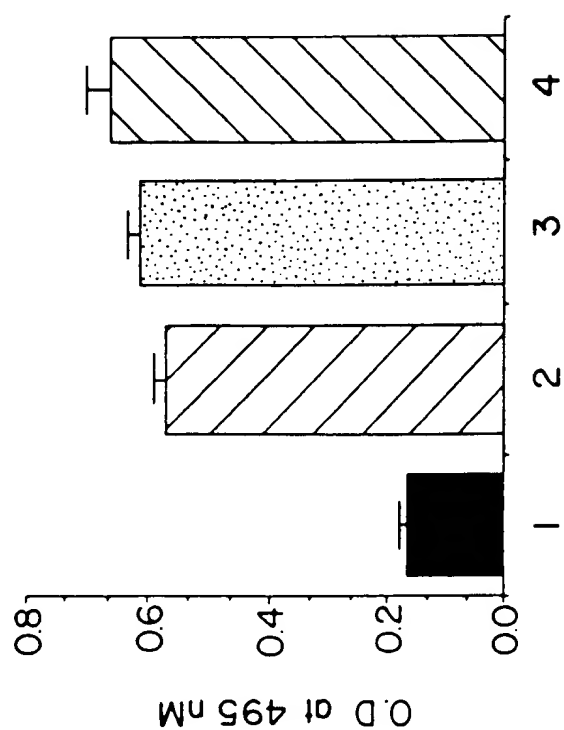
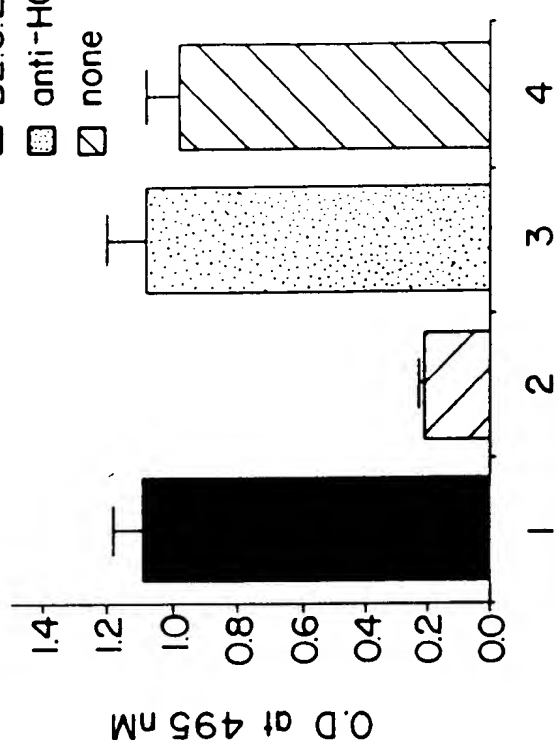


■ A4.6.1
 ▨ B2.6.2
 ▩ anti-HGF
 □ none



BIO-A4.6.1
FIG. 1a

■ A4.6.1
 ▨ B2.6.2
 ▩ anti-HGF
 ▧ none



BIO-B2.6.2
FIG. 1b

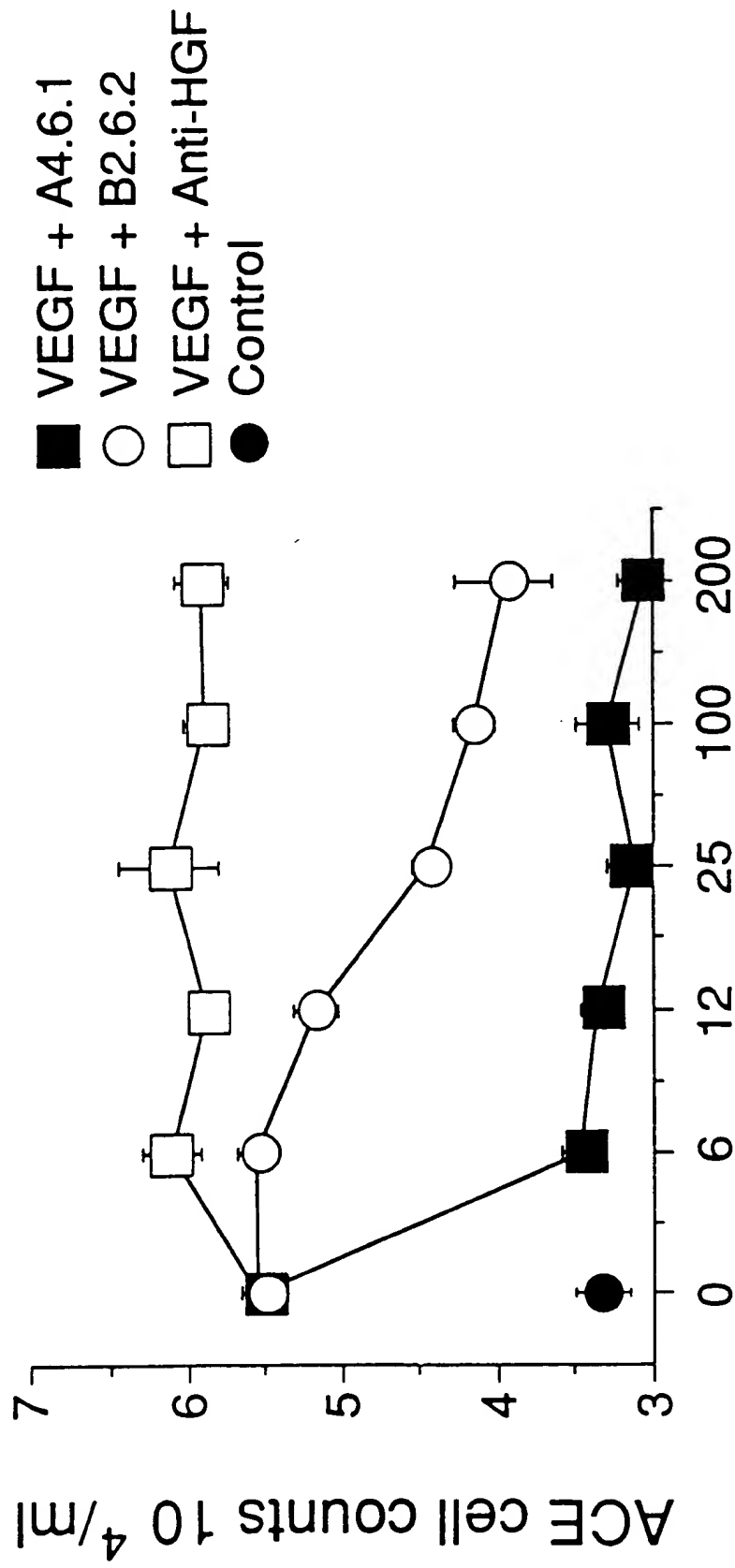


FIG. 2

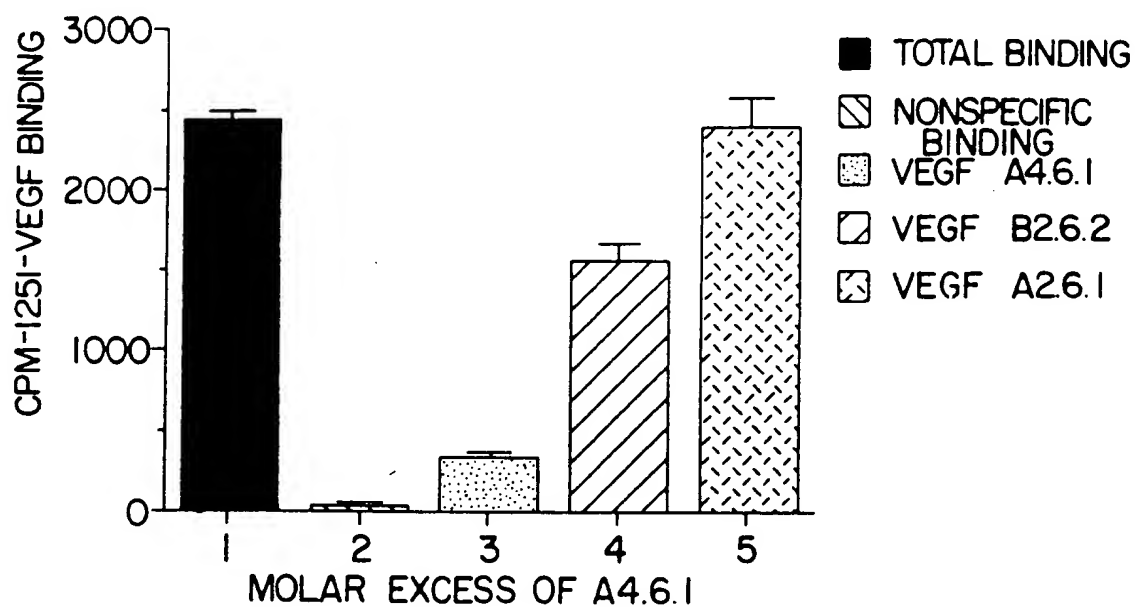


FIG. 3a

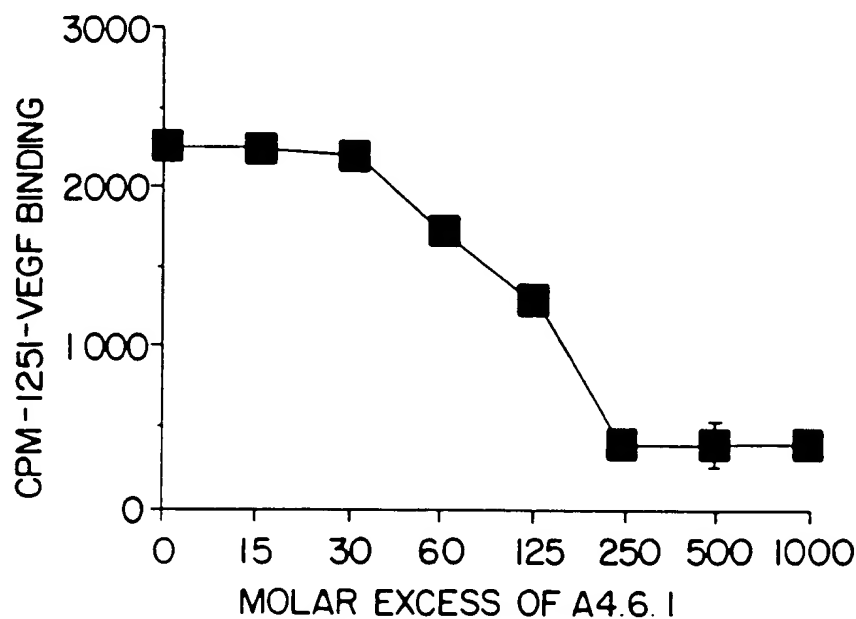


FIG. 3b

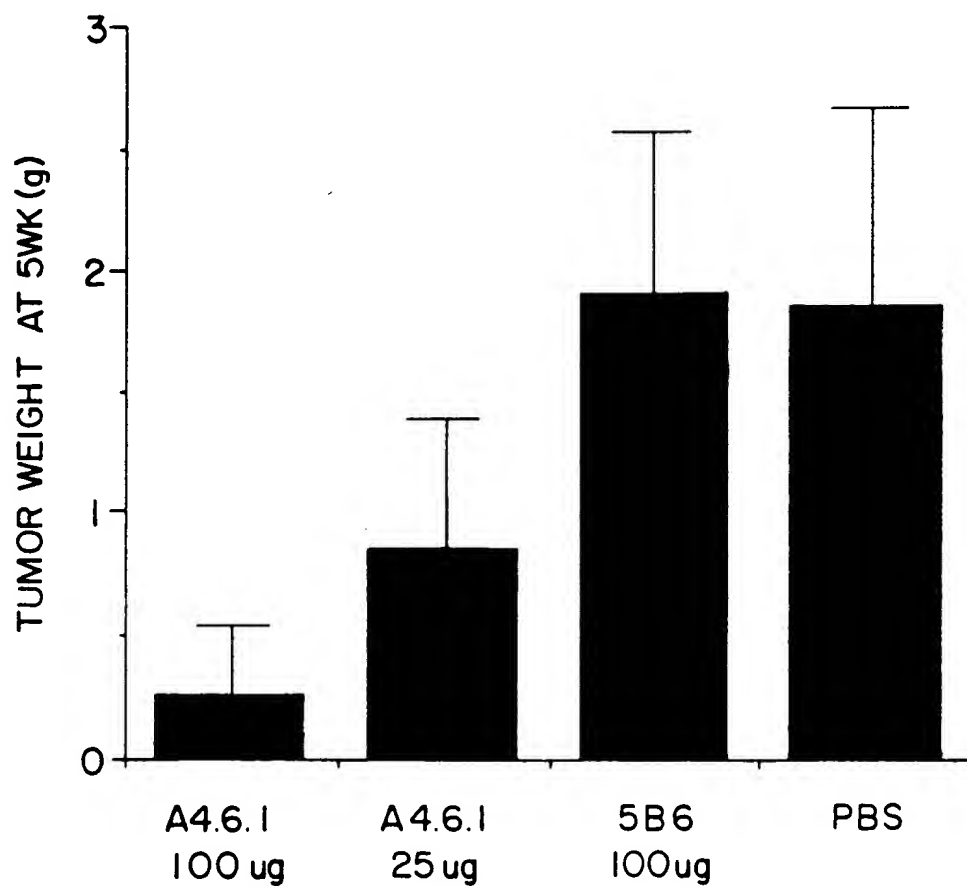


FIG. 5

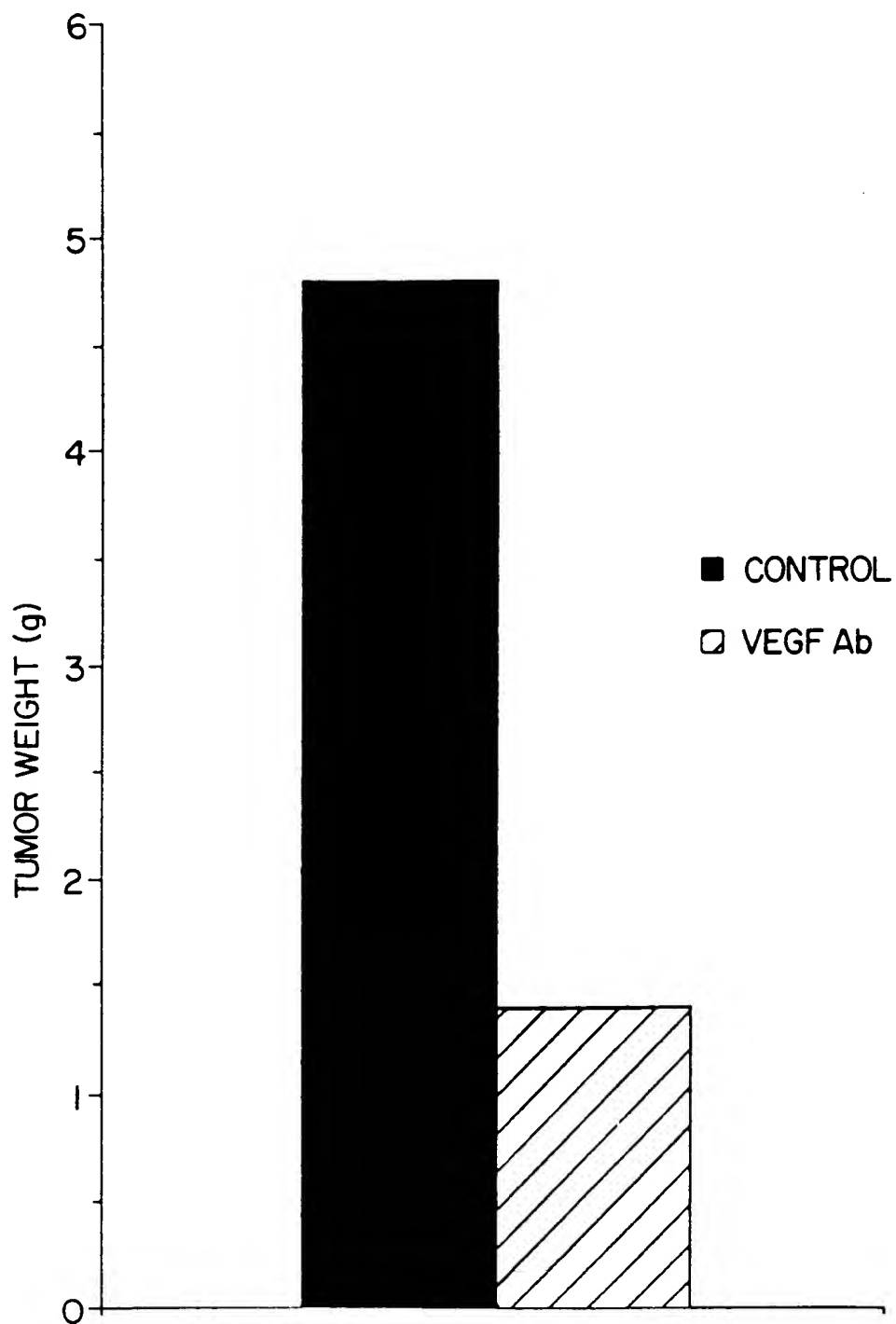


FIG. 6

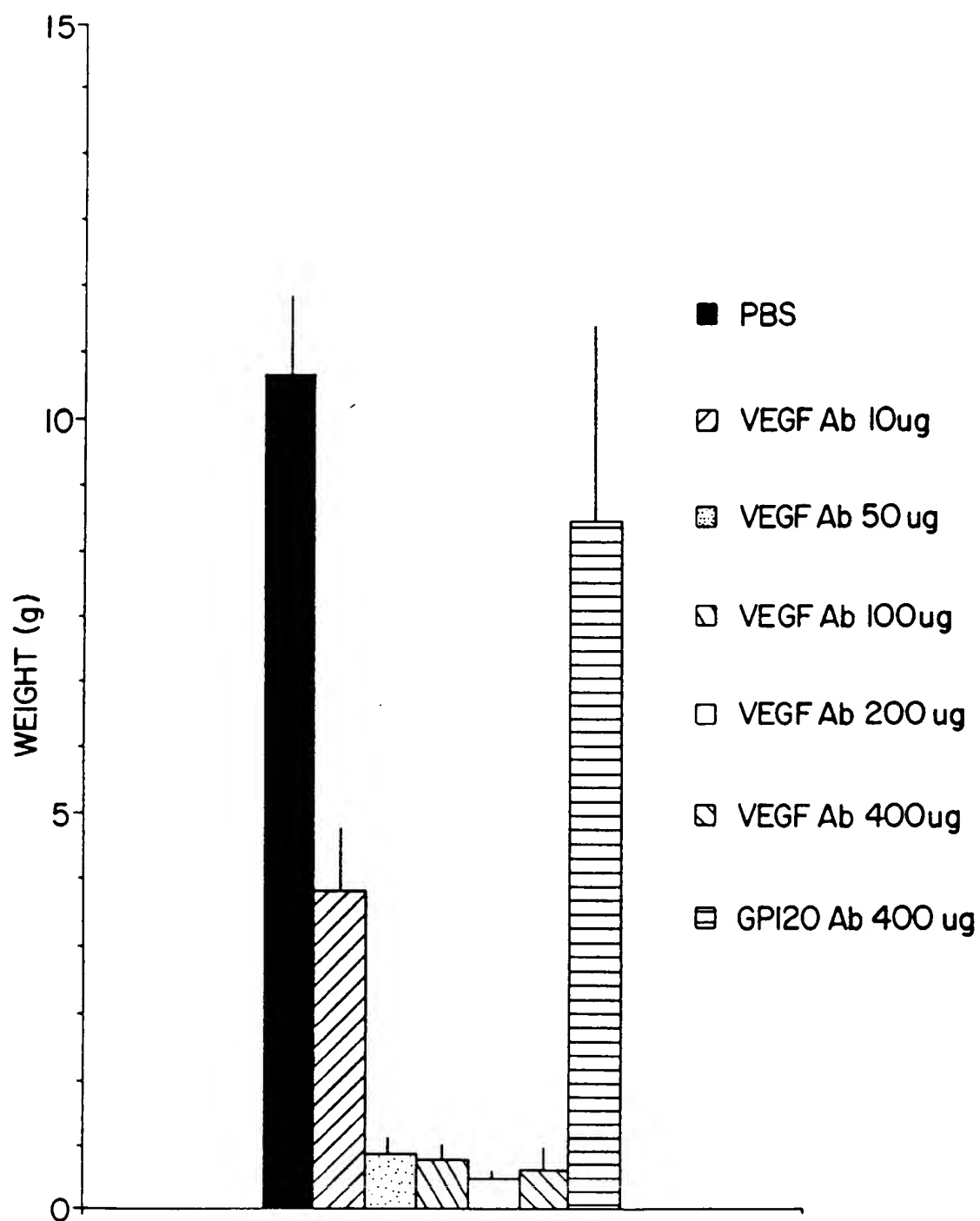


FIG. 7

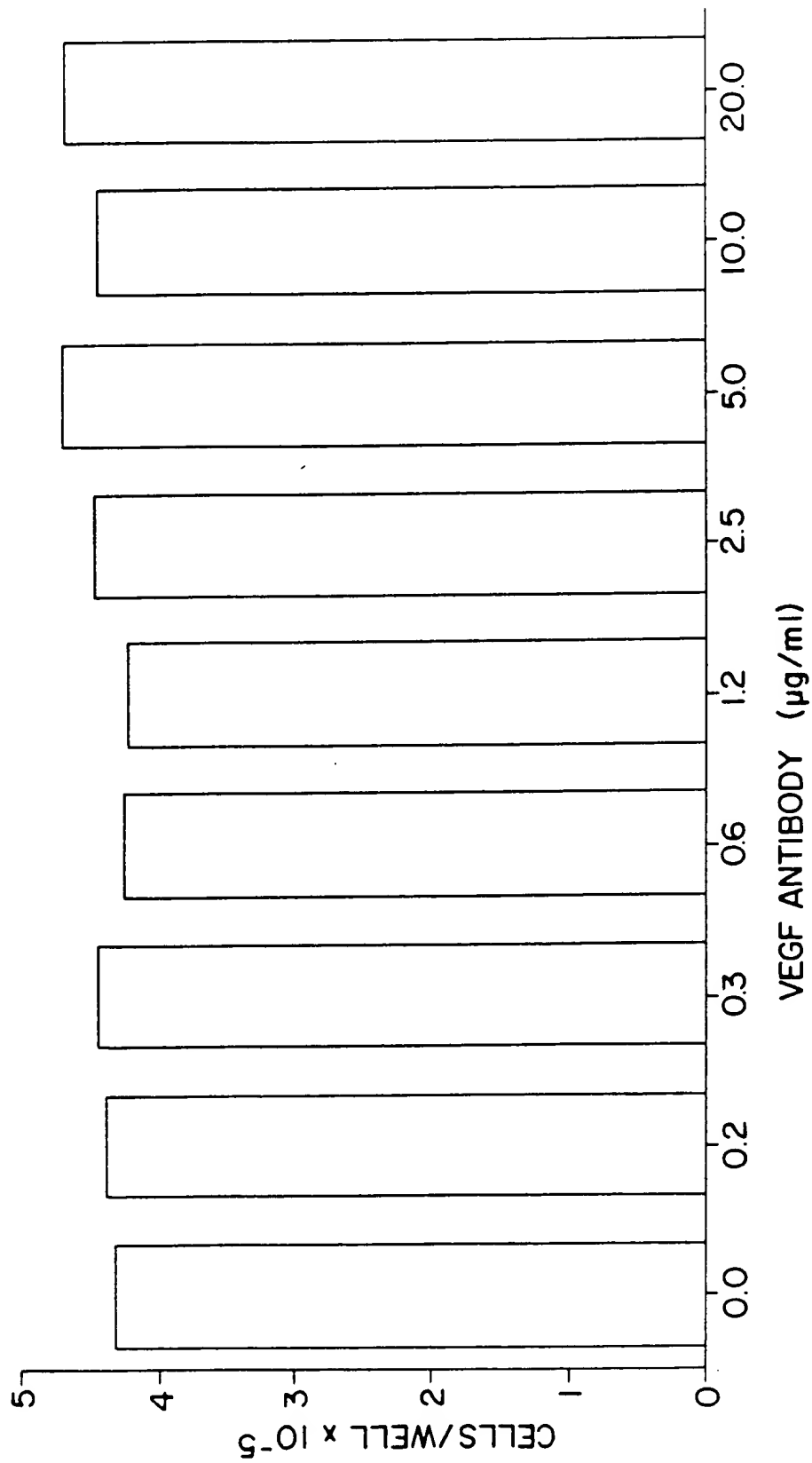


FIG. 8

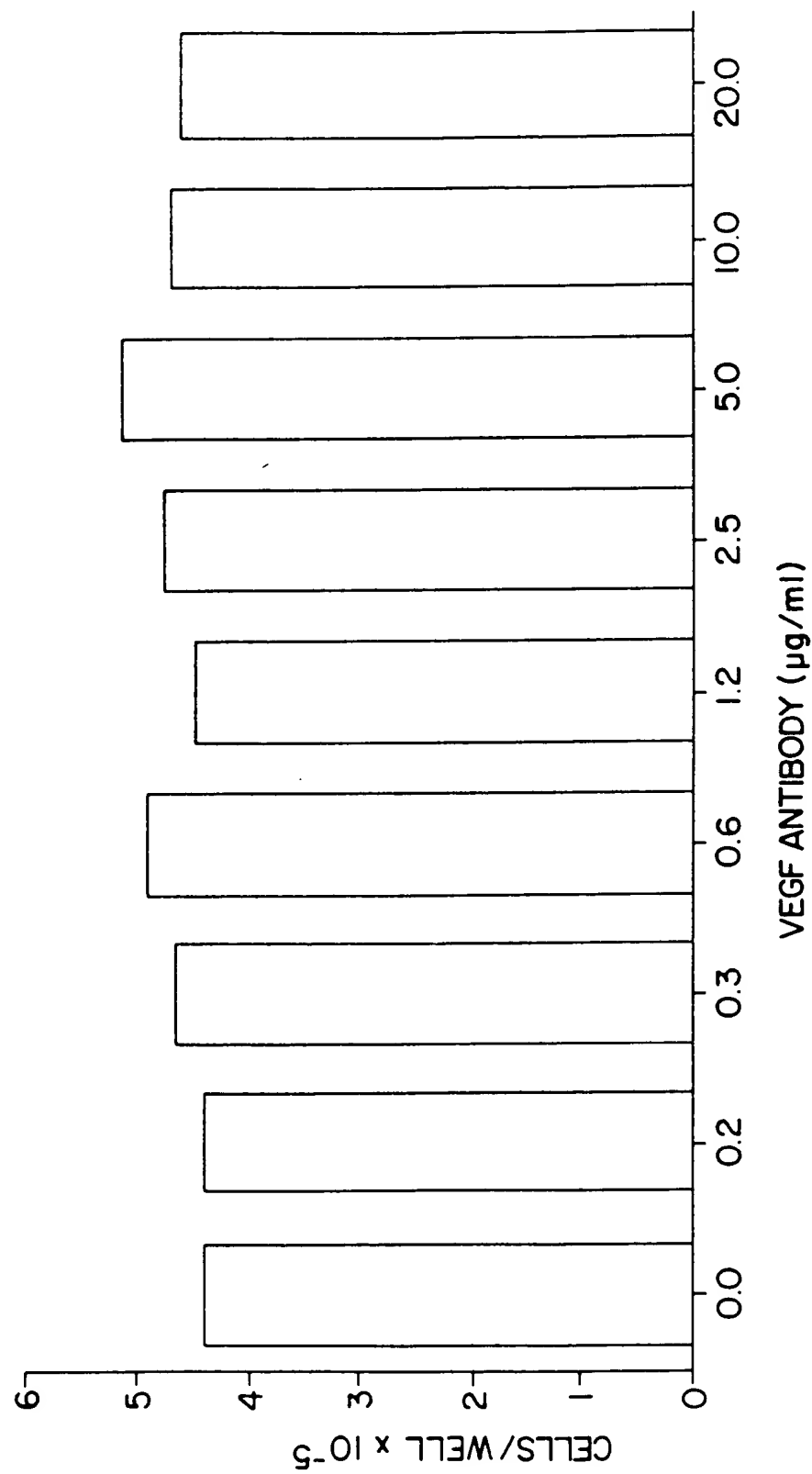


FIG. 9

Sample Type	Sample #	Assay Date	Syn. Fluid	Syn. Fluid + mAB VEGF	% Suppression
Rheumatoid Syn. Fluid	318	5.7.92	5.2 ± 0.2	2.7 ± 0.3	48
	150	5.7.92	7.0 ± 0.3	2.8 ± 0.4	60
	312	5.7.92	6.7 ± 0.4	3.7 ± 0.3	45
	264	5.7.92	6.2 ± 0.4	3.1 ± 0.3	50
	267	5.7.92	5.7 ± 0.6	4.4 ± 0.3	23
	202	5.22.92	10.0 ± 0.5	3.4 ± 0.6	66
	314	5.22.92	7.5 ± 0.3	3.1 ± 0.6	59
	237	5.22.92	6.1 ± 0.5	2.2 ± 0.3	64
	206	5.22.92	6.7 ± 0.5	2.2 ± 0.3	67
	317	5.22.92	5.2 ± 0.3	2.5 ± 0.6	52
Osteoarthritis Syn. Fluid	165	6.2.92	4.0 ± 0.3	2.8 ± 0.4	30
	211	6.2.92	3.4 ± 0.5	3.0 ± 0.2	11.7
	195	6.2.92	3.5 ± 0.2	3.3 ± 0.3	5.7
	122	6.2.92	3.7 ± 0.3	3.2 ± 0.4	13.5
	16	6.2.92	4.1 ± 0.3	3.8 ± 0.5	7.3

Mean % Suppression for RA Fluids 53.4 ± 4.2
Mean % Suppression for OA Fluids 13.6 ± 3.9
Synovial fluids were diluted 1:50.

Controls:

6.2.92	PBS	3.3 ± 0.30
	bFGF 1 µg/ml	3.7 ± 0.38
5.22.92	PBS	1.2 ± 0.38
	bFGF 1 µg/ml	7.8 ± 0.31
5.2.92	PBS	1.3 ± 0.18
	bFGF 1 µg/ml	9.0 ± 0.41

FIG. 10

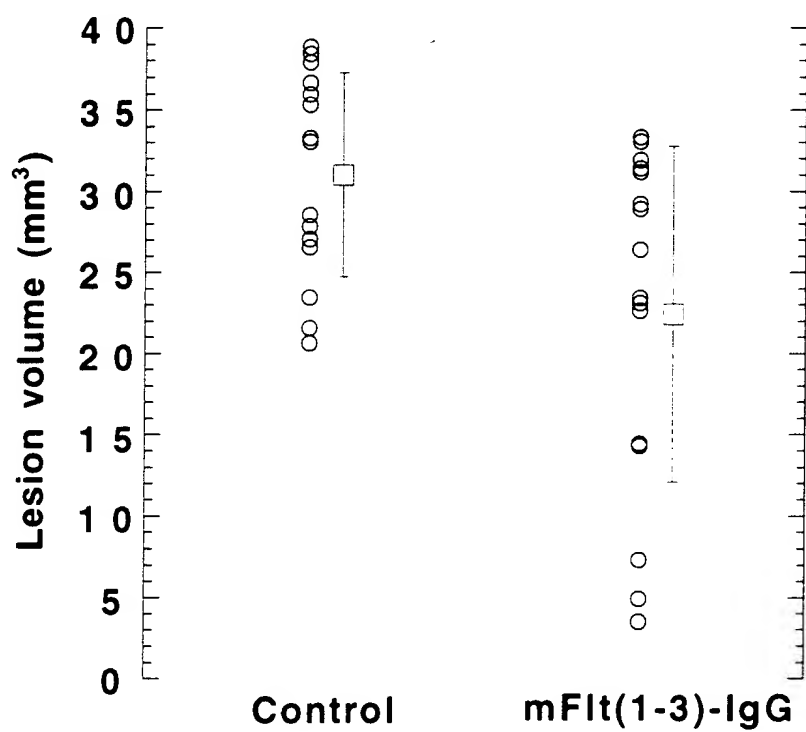


FIG. 11

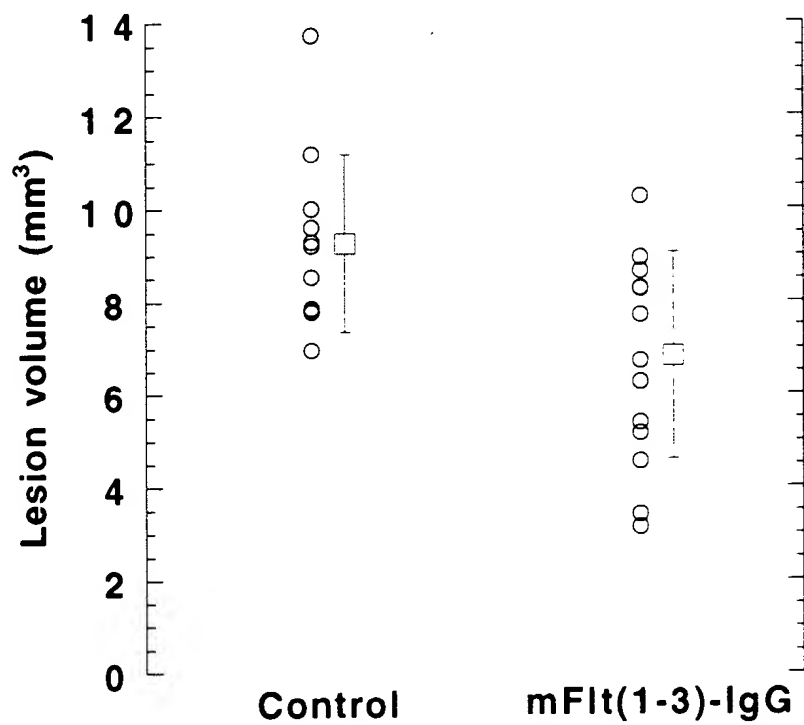


FIG. 13

■ = differences from F-12

10 20 30
F(ab)-12 DIQMTQSPSSLSASVGDRVITITCSASQDISNYLNWYQQ
Y0243-1 DIQMTQSPSSLSASVGDRVITITCSASQDISNYLNWYQQ
Y0238-3 DIQMTQSPSSLSASVGDRVITITCSASQDISNYLNWYQQ
Y0313-1 DIQMTQSPSSLSASVGDRVITITCSASQDISNYLNWYQQ
Y0317 DIQMTQSPSSLSASVGDRVITITCSASQDISNYLNWYQQ

CDR-L1

40 50 60 70
F(ab)-12 KPGKAPKVLIIYFTSSLHSGVPSRFSGSGSGTDFTLTIS
Y0243-1 KPGKAPKVLIIYFTSSLHSGVPSRFSGSGSGTDFTLTIS
Y0238-3 KPGKAPKVLIIYFTSSLHSGVPSRFSGSGSGTDFTLTIS
Y0313-1 KPGKAPKVLIIYFTSSLHSGVPSRFSGSGSGTDFTLTIS
Y0317 KPGKAPKVLIIYFTSSLHSGVPSRFSGSGSGTDFTLTIS

FIG. 14A

CDR-L2

80 90 100
F(ab)-12 SLQPEDFATYYCQQYSTVPWTFGQGTKVEIKRTV
Y0243-1 SLQPEDFATYYCQQYSTVPWTFGQGTKVEIKRTV
Y0238-3 SLQPEDFATYYCQQYSTVPWTFGQGTKVEIKRTV
Y0313-1 SLQPEDFATYYCQQYSTVPWTFGQGTKVEIKRTV
Y0317 SLQPEDFATYYCQQYSTVPWTFGQGTKVEIKRTV

CDR-L3

10 20 30
F(ab)-12 EVQLVESGGGLVQPGGSLRLSCAASGYTFITNYGMNWVR
Y0243-1 EVQLVESGGGLVQPGGSLRLSCAASGYTFITNYGMNWVR
Y0238-3 EVQLVESGGGLVQPGGSLRLSCAASGYTFITNYGMNWVR
Y0313-1 EVQLVESGGGLVQPGGSLRLSCAASGYTFITNYGMNWVR
Y0317 EVQLVESGGGLVQPGGSLRLSCAASGYTFITNYGMNWVR

CDR-H1

40 50 60 70
F(ab)-12 QAPGKGLEWVGWINTYTGEPTYAADFKRRFTFSLDTSKSTA
Y0243-1 QAPGKGLEWVGWINTYTGEPTYAADFKRRFTFSLDTSKSTA
Y0238-3 QAPGKGLEWVGWINTYTGEPTYAADFKRRFTFSLDTSKSTA
Y0313-1 QAPGKGLEWVGWINTYTGEPTYAADFKRRFTFSLDTSKSTA
Y0317 QAPGKGLEWVGWINTYTGEPTYAADFKRRFTFSLDTSKSTA

FIG. 14B

CDR-H2

80 90 100 110 CDR-7
F(ab)-12 YLQMNSLRAEDTAVYYCAKYPHYGSSHWYFDVWGQGTL
Y0243-1 YLQMNSLRAEDTAVYYCAKYPHYGSSHWYFDVWGQGTL
Y0238-3 YLQMNSLRAEDTAVYYCAKYPHYGSSHWYFDVWGQGTL
Y0313-1 YLQMNSLRAEDTAVYYCAKYPHYGSSHWYFDVWGQGTL
Y0317 YLQMNSLRAEDTAVYYCAKYPHYGSSHWYFDVWGQGTL

CDR-H3

[illegible]

F(ab)-12 10 20 30
 Y0192 DIQMTQSPSSLSASVGDRVTTITCSASQDISNYLNWYQQ
 Y0238-3 DIQETQSPSSLSASVGDRVTTITCRANEQLSNYLNWYQQ
 Y0239-19 DIQETQSPSSLSASVGDRVTTITCRANEQLSNYLNWYQQ
 Y0313-2 DIQETQSPSSLSASVGDRVTTITCRANEQLSNYLNWYQQ

CDR-L1

F(ab)-12 KPGKAPKVL⁴⁰LIYFT⁵⁰SSLHSGVPS⁶⁰SRFSGSGSGTDFTLTIS⁷⁰

Y0192 KPGKAPKVL⁴⁰LIYFT⁵⁰SSLHSGVPS⁶⁰SRFSGSGSGTDFTLTIS⁷⁰

Y0238-3 KPGKAPKVL⁴⁰LIYFT⁵⁰SSLHSGVPS⁶⁰SRFSGSGSGTDFTLTIS⁷⁰

Y0239-19 KPGKAPKVL⁴⁰LIYFT⁵⁰SSLHSGVPS⁶⁰SRFSGSGSGTDFTLTIS⁷⁰

Y0313-2 KPGKAPKVL⁴⁰LIYFT⁵⁰SSLHSGVPS⁶⁰SRFSGSGSGTDFTLTIS⁷⁰

CDR-L2

80 90 100

F(ab)-12 SLQPEDFATYYCQQYSTVPWTFGQGTKVEIKRTV

Y0192 SLQPEDFATYYCQQYSTVPWTFGQGTKVEIKRTV

Y0238-3 SLQPEDFATYYCQQYSTVPWTFGQGTKVEIKRTV

Y0239-19 SLQPEDFATYYCQQYSTVPWTFGQGTKVEIKRTV

Y0313-2 SLQPEDFATYYCQQYSTVPWTFGQGTKVEIKRTV

FIG. 15B

F(ab)-12 EVQLVESGGGLVQPGGSLRLSCAASGYTFTNYGMNWVR
Y0192 EVQLVESGGGLVQPGGSLRLSCAASGYTFTNYGMNWVR
Y0238-3 EVQLVESGGGLVQPGGSLRLSCAASGYTFTNYGMNWVR
Y0239-19 EVQLVESGGGLVQPGGSLRLSCAASGYTFTNYGMNWVR
Y0313-2 EVQLVESGGGLVQPGGSLRLSCAASGYTFTNYGMNWVR

CDR-H1

F(ab)-12 40 50 60 70
QAPGKGLEWVGWINTYTGEPYAADF~~KRRFTF~~SLDTSKSTA
Y0192 QAPGKGLEWVGWINTYTGEPYAADF~~KRRFTF~~SLDTSKSTA
Y0238-3 QAPGKGLEWVGWINTYTGEPYAADF~~KRRFTF~~SLDTSKSTA
Y0239-19 QAPGKGLEWVGWINTYTGEPYAADF~~KRRFTF~~SLDTSKSTA
Y0313-2 QAPGKGLEWVGWINTYTGEPYAADF~~KRRFTF~~SLDTSKSTA

	80	90	CDR-H2	100	CDR-7
F(ab)-12	YLQMNSLRAEDTAVYYCAKYPHYYG---	SSHWYFDVWGQGTL			
Y0192	YLQMNSLRAEDTAVYYCAKYPHYYG---	SSHWYFDVWGQGTL			
Y0238-3	YLQMNSLRAEDTAVYYCAKYPHYYG---	SSHWYFDVWGQGTL			
Y0239-19	YLQMNSLRAEDTAVYYCAKYPHYYYVNERK	SSHWYFDVWGQGTL			
Y0313-2	YLQMNSLRAEDTAVYYCAKYPHYYYVNERK	SSHWYFDVWGOGTL			

CDR-H3